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Crystal Structure of the Type III Secretion Chaperone SigE

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Beamline(s): X12C

Introduction: Several gram negative bacterial pathogens have evolved a type III secretion system to deliver virulence effector proteins directly into eukaryotic cells, a process essential for disease [1]. This specialized secretion process requires customized chaperones that are specific for particular effector proteins. Despite the low sequence similarities, most type III secretion chaperones are small (<20 kDa), acidic and dimeric. The crystal structure of the *Salmonella typhimurium* SigD-specific chaperone has been determined at 1.9 Å resolution [2].

Methods and Materials: The crystal structure of intact SigE was solved by MAD phasing using a selenomethionine derivative [3]. All data was collected at 100 K in a cryogenic stream. Three-wavelengths data was collected at the Se absorption peak, edge and high energy remote.

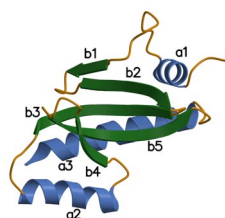
Results: The crystal structure of SigE reveals a novel fold with a-b-b-b-a-b-b-a topology (Fig. 1a). This topology differs significantly from that of SecB, the type II general secretory chaperone [4]. As expected from its acidic pI, SigE has a significant level of negatively charged electrostatic surface area; the charged surface is relatively localized to one face of the molecule and is interspersed with exposed hydrophobic surface (Fig. 1c). Previous work supports the existence of a dimeric form of SigE, *in vivo* and *in vitro*. Accordingly, the asymmetric unit of the SigE crystals contains a homo-dimer in which the two monomers are related by an approximate 2-fold non-crystallographic rotation symmetry. The dimerization interface arises through symmetrical interactions between the region containing the helix-strand motif, $\alpha 2$ -b4, and its dimeric counterpart ($\alpha'1$ -b'4) in such a way that $\alpha 2$ and $\alpha'2$ are approximately parallel (Fig. 1b). The buried area between the dimeric interface is ~2190 Å², predominantly hydrophobic with additional hydrogen bonded interactions by polar side chains.

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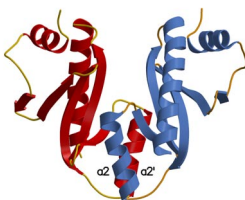
References:

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A.



B.



C.

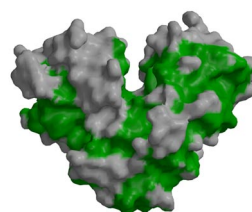


Figure 1. Structure of SigE. A. Ribbon diagram of SigE monomer. B. Ribbon diagram of SigE dimer. C. Surface of SigE (hydrophobic surface in green).